

Package ‘ecorest’

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Title Conducts Analyses Informing Ecosystem Restoration Decisions

Version 2.0.3

Description Three sets of data and functions for informing ecosystem restoration decisions, particularly in the context of the U.S. Army Corps of Engineers. First, model parameters are compiled as a data set and associated metadata for over 300 habitat suitability models developed by the U.S. Fish and Wildlife Service (USFWS 1980, <<https://www.fws.gov/policy-library/870fw1>>). Second, functions for conducting habitat suitability analyses both for the models described above as well as generic user-specified model parameterizations. Third, a suite of decision support tools for conducting cost-effectiveness and incremental cost analyses (Robinson et al. 1995, IWR Report 95-R-1, U.S. Army Corps of Engineers).

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annualizer	<i>Time-averaged restoration project outcomes</i>
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Description

annualizer computes time-averaged quantities based on linear interpolation.

Usage

```
annualizer(timevec, benefits)
```

Arguments

timevec	a numeric vector, matrix, or data frame of time intervals.
benefits	a numeric vector, matrix, or data frame of ecological output values for a single condition (e.g., future with project) to be interpolated.

Value

A time-averaged value over the specified time horizon.

References

Robinson R., Hansen W., and Orth K. 1995. Evaluation of environmental investments procedures manual interim: Cost effectiveness and incremental cost analyses. IWR Report 95-R-1. Institute for Water Resources, U.S. Army Corps of Engineers, Alexandria, Virginia.

Examples

```
#Constant value through time
annualizer(c(0,50), c(100,100))
annualizer(seq(0,50), rep(100,51))

#Simple time series
annualizer(seq(0,50), seq(0,50))

#User-specified time intervals
demo.timevec <- c(0,2,20,50)
demo.ben <- c(0,100,90,80)
annualizer(demo.timevec, demo.ben)
```

BBfinder	<i>Identifies "best buy" actions</i>
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Description

BBfinder this analysis examines the slope of the cost-effectiveness frontier to isolate how unit cost (cost/benefit) increases with increasing environmental benefit. Restoration actions with the lowest slope of unit cost are considered "best buys".

Usage

```
BBfinder(benefit, cost, CE)
```

Arguments

benefit	a vector, matrix, or data frame of restoration benefits. Typically, these are time-averaged ecological outcomes (e.g., average annual habitat units). Often project benefits are best presented as the "lift" associated with a restoration action (i.e., the benefits of an alternative minus the benefits of a "no action" plan).
cost	a vector, matrix, or data frame of restoration costs. Typically, these are monetary costs associated with a given restoration action such as project first cost or annualized economic cost. Notably, these functions are agnostic to units, so costs could also be non-monetary such as lost political capital or social costs of each alternative.
CE	a numeric vector, matrix, or data frame of 0's and 1's indicating whether a plan is cost-effective (1) or non-cost-effective (0). Can be derived from <code>ecorrest::CEfinder</code> .

Value

A list with summaries of all restoration actions as well as best buy plans only.

References

Robinson R., Hansen W., and Orth K. 1995. Evaluation of environmental investments procedures manual interim: Cost effectiveness and incremental cost analyses. IWR Report 95-R-1. Institute for Water Resources, U.S. Army Corps of Engineers, Alexandria, Virginia

Examples

```
#Identify cost-effective actions based on random vectors of benefits and costs
benefit <- runif(50,min=0,max=10)
cost <- runif(50, min=0,max=1000)
CE <- CEfinder(benefit, cost)
BBfinder(benefit, cost, CE)

#Identify cost-effective actions based on a small number of user-specified benefits and costs
restben <- c(0, 10, 5, 20, 20)
restcost <- c(0, 100, 100, 200, 150)
restCE <- CEfinder(restben, restcost)
BBfinder(restben, restcost, restCE)
```

CEfinder

Finds cost-effective frontier

Description

CEfinder returns cost-effectiveness analysis for a particular set of alternatives.

Usage

```
CEfinder(benefit, cost)
```

Arguments

benefit	a vector, matrix, or data frame of restoration benefits. Typically, these are time-averaged ecological outcomes (e.g., average annual habitat units). Often project benefits are best presented as the "lift" associated with a restoration action (i.e., the benefits of an alternative minus the benefits of a "no action" plan).
cost	a vector, matrix, or data frame of restoration costs. Typically, these are monetary costs associated with a given restoration action such as project first cost or annualized economic cost. Notably, these functions are agnostic to units, so costs could also be non-monetary such as lost political capital or social costs of each alternative.

Value

A numeric vector identifying each plan as cost-effective (1) or non-cost-effective (0). The cost-effective actions comprise the Pareto frontier of non-dominated alternatives at a given level of cost or benefit.

References

Robinson R., Hansen W., and Orth K. 1995. Evaluation of environmental investments procedures manual interim: Cost effectiveness and incremental cost analyses. IWR Report 95-R-1. Institute for Water Resources, U.S. Army Corps of Engineers, Alexandria, Virginia

Examples

```
#Identify cost-effective actions based on random vectors of benefits and costs
CEfinder(runif(50,min=0,max=10), runif(50, min=0,max=1000))

#Identify cost-effective actions based on a small number of user-specified benefits and costs
restben <- c(0, 10, 5, 20, 20)
restcost <- c(0, 100, 100, 200, 150)
CEfinder(restben, restcost)
```

CEICAplotter

Plots cost-effectiveness and incremental cost analysis

Description

CEICAplotter Plots Cost-effective Incremental Cost Analysis (CEICA) in *.jpeg format.

Usage

```
CEICAplotter(altnames, benefit, cost, CE, BB, figure.name)
```

Arguments

altnames	a vector of numeric or character values representing unique restoration action identifiers.
benefit	a vector, matrix, or data frame of restoration benefits. Typically, these are time-averaged ecological outcomes (e.g., average annual habitat units). Often project benefits are best presented as the "lift" associated with a restoration action (i.e., the benefits of an alternative minus the benefits of a "no action" plan).
cost	a vector, matrix, or data frame of restoration costs. Typically, these are monetary costs associated with a given restoration action such as project first cost or annualized economic cost. Notably, these functions are agnostic to units, so costs could also be non-monetary such as lost political capital or social costs of each alternative.
CE	a numeric vector, matrix, or data frame of 0's and 1's indicating whether a plan is cost-effective (1) or non-cost-effective (0). Can be derived from ecorest::CEfinder.
BB	a numeric vector, matrix, or data frame of 0's and 1's indicating whether a plan is a best buy (1) or not (0). Can be derived from ecorest::BBfinder.
figure.name	output figure file name structured as "filename.jpeg".

Value

A multi-panel *.jpeg figure summarizing cost-effectiveness and incremental cost analyses.

References

Robinson R., Hansen W., and Orth K. 1995. Evaluation of environmental investments procedures manual interim: Cost effectiveness and incremental cost analyses. IWR Report 95-R-1. Institute for Water Resources, U.S. Army Corps of Engineers, Alexandria, Virginia

Examples

```
#Identify cost-effective actions based on random vectors of benefits and costs
altnames<- paste("Alt",seq(1,50), sep="")
benefit <- runif(50,min=0,max=10)
cost <- runif(50, min=0,max=1000)
CE <- CEfinder(benefit, cost)
BB <- BBfinder(benefit, cost, CE)[[1]][,4]
CEICAplotter(altnames, benefit, cost, CE, BB, tempfile("CEICAexample",fileext=".jpeg"))
```

HSIarimean

Computes Habitat Suitability Index with Arithmetic Mean

Description

HSIarimean uses an arithmetic mean to combine suitability indices into an overarching habitat suitability index. Note that U.S. Army Corps of Engineers users applying the HSIarimean function must have approval from the National Ecosystem Planning Center of Expertise (Eco-PCX) prior to development or application of a new model.

Usage

```
HSIarimean(x)
```

Arguments

x a vector, matrix, or data frame of suitability indices with values ranging from 0 to 1.

Value

A value of habitat quality ranging from 0 to 1 (ignoring NA values).

References

US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.

US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.

US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

Examples

```
#Determine patch quality based on a vector of four suitability indices.
HSIarimean(c(0.1, 0.25, 0.25, 0.25))
```

```
#Determine patch quality based on a vector of suitability indices with an NA.
HSIarimean(c(0.1, 0.25, NA, 0.25))
```

```
#Determine patch quality based on a data frame of suitability indices
x = data.frame(0.1, 0.25, 0.25, 0.25)
colnames(x) = c("var1", "var2", "var3", "var4")
HSIarimean(x)
```

HSIeqtn	<i>Computes Habitat Suitability Index based on Model-Specified Equation</i>
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Description

HSIeqtn computes a habitat suitability index based on equations specified in U.S. Fish and Wildlife Service habitat suitability models contained within `ecorest` via `HSImodels` and `HSImetadata`. Habitat suitability indices represent an overall assessment of habitat quality from combining individual suitability indices for multiple independent variables. The function computes an overall habitat suitability index. Note that models compatible with `HSIeqtn` are certified for use by the U.S. Army Corps of Engineers by the National Ecosystem Planning Center of Expertise (Eco-PCX).

Usage

```
HSIeqtn(HSImodelName, SIV, HSImetadata, exclude = NULL)
```

Arguments

HSImodelname	a character string in quotations that must match an existing model name in <code>HSImetadata</code> .
SIV	a named vector, dataframe, or matrix of suitability index values ranging from 0 to 1 used in the model specified in <code>HSImodelname</code> . Names must correspond exactly with SIV names (e.g., "SIV1", "SIV1B", "SIV2", etc.)
HSImetadata	a data frame of HSI model metadata within the <code>ecorest</code> package.

exclude a vector of character strings specifying components to be excluded from calculations. Non-NULL 'exclude' inputs are only applicable to models that explicitly provide instructions for use in the note column of HSI metadata.

Value

A numeric value between zero and one representing the composite habitat suitability index score.

References

US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.

US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.

US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

Examples

```
#Compute patch quality for the Barred Owl model (no components)
#Allen A.W. 1982. Habitat Suitability Index Models: Barred owl. FWS/OBS 82/10.143.
#U.S. Fish and Wildlife Service. https://pubs.er.usgs.gov/publication/fwsobs82_10_143.
#Suitability indices relate to density of large trees (SIV1),
#mean diameter of overstory trees (SIV2), and percent canopy cover of overstory (SIV3).
HSIeqtn("barredowl", c("SIV1" = 1, "SIV2" = 1, "SIV3" = 1),
HSImetadata) #c(1,1,1) should result in 1.00
HSIeqtn("barredowl", c("SIV1" = 0.5, "SIV2" = 1, "SIV3" = 1),
HSImetadata) #c(0.5,1,1) should result in 0.707
HSIeqtn("barredowl", c("SIV1" = 0, "SIV2" = 1, "SIV3" = 1),
HSImetadata) #c(0,1,1) should result in 0.00

#Compute patch quality for the Juvenile Alewife model (two components)
#Pardue, G.B. 1983. Habitat Suitability index models: alewife and blueback herring.
#U.S. Dept. Int. Fish Wildl. Serv. FWS/OBS-82/10.58. 22pp.
#Suitability indices relate to zooplankton density, salinity, and water temperature
HSIeqtn("alewifeJuv", c("SIV1" = NA, "SIV2" = NA,
" SIV3" = 1, "SIV4" = 1, "SIV5" = 1), HSI metadata) #should result in 1.00
HSIeqtn("alewifeJuv", c("SIV1" = NA, "SIV2" = NA,
" SIV3" = 0.5, "SIV4" = 1, "SIV5" = 1), HSI metadata) # should result in 0.50
HSIeqtn("alewifeJuv", c("SIV1" = NA, "SIV2" = NA,
" SIV3" = 0, "SIV4" = 1, "SIV5" = 1), HSI metadata) # should result in 0.00

#Compute patch quality for Cutthroat trout model for lacustrine habitats (7 components)
#with spawning and lacustrine habitat and with only lacustrine habitat (i.e.,
#embryo component is excluded).
#Hickman, T., and R.F. Raleigh. 1982. Habitat suitability index models:
#Cutthroat trout. U.S.D.I. Fish and Wildlife Service. FWS/OBS-82/10.5. 38 pp.
#Suitability indices relate to temperature during the warmest period of the year,
#maximum temperature during embryo development, minimum dissolved oxygen during
#the late growing season, average velocity over spawning areas, average size
#of substrate in spawning areas, annual maximal or minimal pH, and percent fines
```

```

#in the spawning area.
#should result in 1
HSIeqtn("cutthroatLacGenLtoe15C", c("SIV1" = 1, "SIV2" = 1, "SIV3" = 1,
                                     "SIV5" = 1, "SIV7" = 1, "SIV13" = 1,
                                     "SIV16" = 1), HSImetadata)

#should result in 0
HSIeqtn("cutthroatLacGenLtoe15C", c("SIV1" = 0.5, "SIV2" = 1, "SIV3" = 0.5,
                                     "SIV5" = 0, "SIV7" = 1, "SIV13" = 1,
                                     "SIV16" = 1), HSImetadata)

#should result in 0.63
HSIeqtn("cutthroatLacGenLtoe15C", c("SIV1" = 1, "SIV2" = NA, "SIV3" = 0.5,
                                     "SIV5" = NA, "SIV7" = NA, "SIV13" = 0.5,
                                     "SIV16" = NA), HSImetadata, exclude=c("CE"))

```

HSIgeomean

Habitat Suitability Index with Geometric Mean

Description

HSIgeomean uses a geometric mean to combine suitability indices into an overarching habitat suitability index. Note that U.S. Army Corps of Engineers users applying the HSIgeomean function must have approval from the National Ecosystem Planning Center of Expertise (Eco-PCX) prior to application of a new model.

Usage

```
HSIgeomean(x)
```

Arguments

x a vector, matrix, or data frame of suitability indices with values ranging from 0 to 1.

Value

A value of habitat quality ranging from 0 to 1 (ignoring NA values).

References

- US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.
- US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.
- US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

Examples

```
#Determine patch quality based on a vector of four suitability indices.
HSIgeomean(c(0.25, 0.25, 0.25, 0.25))

#Determine patch quality based on a vector of suitability indices with an NA.
HSIgeomean(c(0.25, 0.25, NA, 0.25))

#Determine patch quality based on a vector of suitability indices with a zero-value.
HSIgeomean(c(0.25, 0.25, 0.0, 0.25))

#Determine patch quality based on a dataframe of suitability indices
x = data.frame(0.25, 0.25, 0.25, 0.25)
colnames(x) = c("var1", "var2", "var3", "var4")
HSIgeomean(x)
```

HSImetadata

Habitat suitability index (HSI) model metadata

Description

Metadata for 351 U.S. Fish and Wildlife Service Habitat suitability index (HSI) models

Usage

HSImetadata

Format

A data frame with 351 rows and 85 variables:

model Model name
submodel Model specifications
species Scientific nomenclature of modeled taxa
geography Geographic range of organism
ecosystem Type of habitat
documentation Citation of original model
note Conditions under which model may be applied
website Link to individual model source
SIV1 Suitability index values for each organism specific condition
SIV1B Suitability index values for each organism specific condition
SIV2 Suitability index values for each organism specific condition
SIV2B Suitability index values for each organism specific condition
SIV3 Suitability index values for each organism specific condition

SIV3B Suitability index values for each organism specific condition
SIV4 Suitability index values for each organism specific condition
SIV4B Suitability index values for each organism specific condition
SIV5 Suitability index values for each organism specific condition
SIV5B Suitability index values for each organism specific condition
SIV6 Suitability index values for each organism specific condition
SIV6B Suitability index values for each organism specific condition
SIV7 Suitability index values for each organism specific condition
SIV7B Suitability index values for each organism specific condition
SIV8 Suitability index values for each organism specific condition
SIV8B Suitability index values for each organism specific condition
SIV9 Suitability index values for each organism specific condition
SIV10 Suitability index values for each organism specific condition
SIV11 Suitability index values for each organism specific condition
SIV12 Suitability index values for each organism specific condition
SIV13 Suitability index values for each organism specific condition
SIV14 Suitability index values for each organism specific condition
SIV15 Suitability index values for each organism specific condition
SIV15B Suitability index values for each organism specific condition
SIV16 Suitability index values for each organism specific condition
SIV16B Suitability index values for each organism specific condition
SIV17 Suitability index values for each organism specific condition
SIV18 Suitability index values for each organism specific condition
SIV19 Suitability index values for each organism specific condition
SIV20 Suitability index values for each organism specific condition
SIV21 Suitability index values for each organism specific condition
SIV22 Suitability index values for each organism specific condition
CF Food component equation
CRF Food/reproduction component equation
CRN Roosting-nesting component equation
CC Cover component equation
CCRO Cover roosting component equation
CCRF Cover-reproduction-food component equation
CCF Cover-food component equation
CCSF Cover-food shrub component equation
CCHF Cover-food herbaceous component equation
CWF Winter food component

CSF Summer food component
CFF Fall food component
CW Water component
CCB Cover breeding component
CB Brood component
CN Nest component
CNBC Nest-brood cover component
CCN Cover nesting component
CP Pair habitat component
CWQ Water quality component
CR Reproduction component
CCR Cover reproduction component
CD Disturbance component
COT Other component
CL Larval component
CEL Embryo and larval component
CE Embryo component
CJ Juvenile component
CFr Fry component
CS Spawning component
CA Adult component
CI Island component
CIN Interspersion component
CNI Non-island component
CWFC Winter cover food component
CFBS Summer food brood component
CFSWF Fall spring winter food component
CSPF Spring food component
CWC Winter cover component
CCFS Fall to spring cover component
CSS Substrate-suspended solids component
CT Topography component
CTe Temperature component
CJA Juvenile adult component
Eqtn HSI overarching model equation in R syntax

Source

<https://pubs.usgs.gov/>

HSImin*Habitat Suitability Index with Limiting Factors*

Description

HSImin uses the minimum of given suitability indices to calculate an overarching habitat suitability index. Note that U.S. Army Corps of Engineers users applying the HSImin function must have approval from the National Ecosystem Planning Center of Expertise (Eco-PCX) prior to application of a new model.

Usage

```
HSImin(x)
```

Arguments

x a vector, matrix, or data frame of suitability indices ranging from 0 to 1.

Value

A value of habitat quality ranging from 0 to 1 (ignoring NA values).

References

US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.

US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.

US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

Examples

```
#Determine patch quality based on a vector of four suitability indices.  
HSImin(c(0.1, 0.25, 0.25, 0.25))
```

```
#Determine patch quality based on a vector of suitability indices with an NA.  
HSImin(c(0.1, 0.25, NA, 0.25))
```

```
#Determine patch quality based on a data frame of suitability indices  
x = data.frame(0.1, 0.25, 0.25, 0.25)  
colnames(x) = c("var1", "var2", "var3", "var4")  
HSImin(x)
```

HSImodels

Habitat suitability index (HSI) models

Description

This list of data frames contains 351 U.S. Fish and Wildlife Service Habitat suitability index (HSI) models. Please note that some of the original HSI documents provide little reference data for constructing suitability curves; hence, some suitability curves are estimated using the authors' best judgement. Users should always cross-reference results with the original documentation.

Usage

HSImodels

Format

An object of class `list` of length 349.

Details

@format A list with 351 data frames each containing an HSI model with multiple independent variables and associated habitat suitability indices (a 0 to 1 value). Data represent break points in curves with linear extrapolation between. Categorical input variables are coded as letters.

variable1 independent variable for assessing habitat suitability

SIV1 suitability index value relative to variable1

... additional variables and suitability indices

Source

<https://pubs.usgs.gov/>

HSIplotter

Plots habitat suitability index curves

Description

HSIplotter plots all suitability curves. Note that U.S. Army Corps of Engineers users must have approval from the National Ecosystem Planning Center of Expertise (Eco-PCX) prior to application of a new model.

Usage

HSIplotter(SI, figure.name)

Arguments

SI list, matrix, or data frame of suitability curves ordered as parameter breakpoints and associated suitability indices for each parameter with appropriate column names. Models containing both categorical and continuous parameters must be entered as a dataframe.

figure.name output figure file name structured as "filename.jpeg".

Value

A multi-panel *.jpeg figure showing all suitability curves.

References

US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.

US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.

US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

Examples

```
#Build and define a matrix of the Barred Owl suitability curves
#Allen A.W. 1982. Habitat Suitability Index Models: Barred owl. FWS/OBS 82/10.143.
#U.S. Fish and Wildlife Service. https://pubs.er.usgs.gov/publication/fwso82_10_143.
var1 <- cbind(c(0,2,4,NA), c(0.1,1,1,NA)) #Number of trees > 51cm diameter per 0.4 ha plot
var2 <- cbind(c(0,5,20,NA), c(0,0,1,NA)) #Mean diameter of overstory trees
var3 <- cbind(c(0,20,60,100), c(0,0,1,1)) #Percent canopy cover of overstory trees
barredowl <- cbind(var1, var2, var3)
colnames(barredowl)<- c("tree.num", "tree.num.SIV",
  "avg.dbh.in", "avg.dbh.SIV", "can.cov", "can.cov.SIV")

#Create suitability curve summary plot
HSIplotter(barredowl, tempfile("BarredOwl",fileext=".jpeg"))

#Build and define a matrix of the alewifeSAEL curves
#Pardue, GB. 1983. Habitat suitability index models: alewife and blueback herring.
#U.S. Dept. Int. Fish Wildl. Serv. FWS/OBS-82/10.58. 22pp.
var1 = data.frame(subs.type.class = c("a", "b", "c", NA, NA, NA),
  subs.type.class.SIV = c(1, 0.5, 0.1, NA, NA, NA))
var2 = data.frame("avg.daily.wtr.temp.spwn.C" = c(5, 10, 15, 20, 27, 30),
  "avg.daily.wtr.temp.spwn.C.SIV" = c(0, 0, 1, 1, 0, 0))
alewifeSAEL = data.frame(var1, var2)

#Create suitability curve summary plot
HSIplotter(alewifeSAEL, tempfile("AlewifeSAEL",fileext=".jpeg"))

#Use HSIplotter with inputs from HSImodels
HSIplotter(HSImodels$barredowl, tempfile("BarredOwl",fileext=".jpeg"))
```

 HSIwarimean

Habitat Suitability Index with a Weighted Arithmetic Mean

Description

HSIwarimean uses a weighted arithmetic mean to combine suitability indices into an overarching habitat suitability index. Note that U.S. Army Corps of Engineers users applying the HSIwarimean function must have approval from the National Ecosystem Planning Center of Expertise (Eco-PCX) prior to application of a new model or weighting system.

Usage

```
HSIwarimean(x, w)
```

Arguments

x is a vector, matrix, or data frame of suitability indices ranging from 0 to 1.
w is a vector, matrix, or data frame of weights ranging from 0 to 1 that must sum to one.

Value

A value of habitat quality ranging from 0 to 1 (ignoring NA values).

References

US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.
 US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.
 US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

Examples

```
#Determine patch quality based on a vector of four, equal-weight suitability indices.
HSIwarimean(c(1, 0, 0, 0), c(0.25, 0.25, 0.25, 0.25))

#Determine patch quality based on a vector of four, unequal-weight suitability indices.
HSIwarimean(c(1, 0, 0, 0), c(1, 0, 0, 0))

#Determine patch quality based on a vector of four, unequal-weight suitability indices.
HSIwarimean(c(1, 0, 0, 0), c(0, 1, 0, 0))

#Determine patch quality based on a data frame of four, unequal-weight suitability indices
df = data.frame(x = c(0.25, 0.5, 0.5, 0.5), w = c(0.25, 0.2, 0.5, 0.05))
```

```
HSIwarimean(df$x, df$w)
```

 HUcalc

Computes Habitat Quality, Quantity, and Units

Description

HUcalc computes habitat units given a set of suitability indices, a habitat suitability index equation, and habitat quantity. Note that U.S. Army Corps of Engineers users applying HUcalc must have approval from the National Ecosystem Planning Center of Expertise (Eco-PCX) prior to application of a new model.

Usage

```
HUcalc(SI.out, habitat.quantity, HSIfunc, ...)
```

Arguments

SI.out	is a vector, matrix, or data frame of application-specific suitability indices between 0 and 1, which can be produced from SICalc.
habitat.quantity	is a numeric value representing habitat size associated with these suitability indices (i.e., length, area, or volume).
HSIfunc	is a function used to combine suitability indices into a composite habitat suitability index (HSI score) (e.g., ecorest functions like HSIarimean or HSIgeomean or functions outside ecorest like max or mean)
...	optional arguments to HSIfunc.

Value

A vector of habitat quality, habitat quantity, and index units (quantity times quality).

References

- US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.
- US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.
- US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

Examples

```
#Summarize habitat outcomes based on a vector of two suitability indices
#using multiple combination equations.
HUcalc(c(0.1,1), 100, HSIarimean)
HUcalc(c(0.1,1), 100, HSIgeomean)
HUcalc(c(0.1,1), 100, HSImin)
HUcalc(c(0.1,1), 100, HSIwarimean, c(1,0))
HUcalc(c(0.1,1), 100, HSIwarimean, c(0,1))

#HSIfunc can also represent functions outside of the ecorest package
HUcalc(c(0.1,1), 100, mean)
HUcalc(c(0.1,1), 100, max)

#Summarize habitat outcomes based on a data frame of three suitability indices
SI.out = data.frame(SI = c(0.1, 0.4, 0.5))
HUcalc(SI.out, 10, HSImin)
```

Sicalc

Computes Suitability Indices

Description

Sicalc computes suitability indices given a set of suitability curves and project-specific inputs. Suitability indices may be computed based on either linear interpolation (for continuous variables) or a lookup method (for categorical variables).

Usage

```
Sicalc(SI, input.proj)
```

Arguments

SI	matrix or dataframe of suitability curves ordered as parameter breakpoints and associated suitability indices for each parameter. Suitability curves that contain both continuous and categorical variables should be formatted as a dataframe rather than a matrix.
input.proj	numeric or categorical vector of application-specific input parameters associated with the suitability curve data from SI. Note that users should enter NA for excluded variables in HSImodels.

Value

A named vector of suitability index values ranging from 0 to 1 that match given user inputs.

References

- US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.
- US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.
- US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

Examples

```
#Build and define a matrix of the Barred Owl suitability curves
#Allen A.W. 1982. Habitat Suitability Index Models: Barred owl. FWS/OBS 82/10.143.
#U.S. Fish and Wildlife Service. https://pubs.er.usgs.gov/publication/fwsobs82\_10\_143.
var1 <- cbind(c(0,2,4,NA), c(0.1,1,1,NA)) #Number of trees > 51cm diameter per 0.4 ha plot
var2 <- cbind(c(0,13,51,NA), c(0,0,1,NA)) #Mean diameter of overstory trees
var3 <- cbind(c(0,20,60,100), c(0,0,1,NA)) #Percent canopy cover of overstory trees
barredowl <- cbind(var1, var2, var3)
colnames(barredowl)<- c("tree.num", "tree.num.SIV",
  "avg.dbh.cm", "avg.dbh.SIV", "can.cov", "can.cov.SIV")

#Set user input variables (in cm) that should return (1, 0, 0)
input.demo1 <- c(2, 0, 0)
Sicalc(barredowl, input.demo1)

#Set user input variables (in cm) that should return (1, 1, 1)
input.demo2 <- c(4, 51, 60)
Sicalc(barredowl, input.demo2)

#Set user input variables (in cm) that should return (1, 1, 0.5)
input.demo3 <- c(4, 51, 40)
Sicalc(barredowl, input.demo3)

#Suitability curves may also be drawn from HSIModels (data within ecoest)
#Import Barred Owl suitability curves with HSIModels$barredowl
#The input examples are repeated from above

#Set user input variables that should return (1, 0, 0)
Sicalc(HSIModels$barredowl, input.demo1)

#Set user input variables that should return (1, 1, 1)
Sicalc(HSIModels$barredowl, input.demo2)

#Set user input variables that should return (1, 1, 0.5)
Sicalc(HSIModels$barredowl, input.demo3)

#Import juvenile Alewife suitability curves with HSIModels$alewifeJuv
#Demonstrate how to enter NA for excluded variables in HSIModels
#Pardue, GB. 1983. Habitat suitability index models: alewife and blueback herring.
#U.S. Dept. Int. Fish Wildl. Serv. FWS/OBS-82/10.58. 22pp.
```

```
#Set user variables that should return (NA, NA, 1, 1, 0)
input.demo4 <- c(NA, NA, 125, 5, 5)
Sicalc(HSImodels$salewifeJuv, input.demo4)

#Demonstrate use with out of range continuous inputs
input.demo5 <- c(NA, NA, 140, 5, 5)
Sicalc(HSImodels$salewifeJuv, input.demo5)
```

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